## SEQUENCE LISTING

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<110> Hitoshi, Yasumichi
      Anderson, David
      Rigel Pharmaceuticals, Inc.
<120> Cell Cycle Targets and Peptides
<130> 021044-002430PC
<140> US 10/531,492
<141> 2005-04-15
<150> US 60/422,912
<151> 2002-10-30
<150> US 60/460,845
<151> 2003-04-04
<150> WO PCT/US03/34669
<151> 2003-10-30
<160> 58
<170> PatentIn Ver. 2.1
<210> 1
<211> 20
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      peptide 88, peptide 35/88
<400> 1
Arg Leu Arg Arg Ile Cys Ser Gly Ile Leu Leu Ile Arg Arg Ile Leu
                  5
  1
                                      10
Gly Ile Phe Val
<210> 2
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<223> Description of Artificial Sequence: C-terminus
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<400> 2
Arg Pro Val Arg
<210> 3
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     vector-derived sequence
<400> 3
Arg Pro Val Arg Pro
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<400> 4
Thr Ser Gly Leu Leu Lys Leu Val Gln Ala Lys Arg Lys Cys Cys Ile
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                  5
                                      10
                                                          15
Ser
<210> 5
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<223> Description of Artificial Sequence:peptide 40
Arg Trp Asp Pro Thr Arg Leu Leu Arg Phe Arg Phe Leu Arg Met Leu
                                      10
Val Arg Arg Ser
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<210> 6
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<400> 6
Gly Arg Gly Cys Ile Phe Arg Trp Arg Gly Leu Arg Gly Met Met
Arg Leu Phe Lys
<210> 7
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<220>
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      fused to N-terminus, K7, lys7
<400> 7
Lys Lys Lys Lys Lys Lys
<210> 8
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<223> Description of Artificial Sequence: nucleotide
      sequence encoding peptide 35/88
<400> 8
cggctccgga gaatatgtag cggcattctg ctcatccgta ggatattggg cattttcgtt 60
aggcccgtga ggccctaa
<210> 9
<211> 53
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      sequence encoding peptide 38
actagtgggt tgctgaagct ggtgcaggct aagcgtaagt gttgtattag tta
                                                             53
<210> 10
<211> 78
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<400> 10
cgttgggatc cgacgcgatt gctgcgattt cggttcctcc ggatgctagt gaggcggagt 60
aggcccgtga ggccctaa
<210> 11
<211> 63
<212> DNA
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<223> Description of Artificial Sequence:nucleotide
      sequence encoding peptide 41
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ggaaggggat gtatctttcg atggaggaga ggcctgcggg gaatgatgag actatttaag 60
<210> 12
<211> 8
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<221> MOD RES
<222> (1)..(3)
<223> Xaa = large hydrophobic amino acid selected from
      the group Leu, Ile, Phe, Met, Tyr or Trp, wherein
      at least one is Leu or Ile, Xaa at position 3 may
      be present or absent
<220>
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<222> (4)..(6)
<223> Xaa = large hydrophobic amino acid selected from
      the group Leu, Ile, Phe, Met, Tyr or Trp, wherein
      at least one is Leu or Ile, Xaa at position 6 may
      be present or absent
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<221> MOD_RES
<222> (7)..(8)
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      the group Leu, Ile, Phe, Met, Tyr or Trp, wherein
      at least one is Leu or Ile
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<400> 12
Xaa Xaa Xaa Xaa Xaa Xaa Xaa
                  5
<210> 13
<211> 6
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:linker
<400> 13
Glu Glu Ala Ala Lys Ala
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  1
<210> 14
<211> 37
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: biotinylated
      peptide 40 fused to C-terminus of GFP
<220>
<221> MOD_RES
<222> (1)
<223> Xaa = biotinylated Gly
<400> 14
Xaa Met Asp Glu Leu Tyr Lys Glu Glu Ala Ala Lys Ala Arg Trp Asp
Pro Thr Arg Leu Leu Arg Phe Arg Phe Leu Arg Met Leu Val Arg Arg
                                 25
Ser Arg Pro Val Arg
         35
<210> 15
<211> 37
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      biotinylated alanine mutant of peptide 40
<220>
<221> MOD_RES
<222> (1)
<223> Xaa = biotinylated Gly
<400> 15
Xaa Met Asp Glu Leu Tyr Lys Glu Glu Ala Ala Lys Ala Arg Trp Asp
Pro Thr Arg Ala Leu Arg Ala Arg Phe Ala Arg Ala Leu Val Arg Arg
             20
Ser Arg Pro Val Arg
         35
<210> 16
<211> 33
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence:peptide 41
      fused to C-terminus of GFP
<400> 16
Gly Met Asp Glu Leu Tyr Lys Glu Glu Ala Ala Lys Ala Gly Arg Gly
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Cys Ile Phe Arg Trp Arg Arg Gly Leu Arg Gly Met Met Arg Leu Phe
             20
                                  25
Lys
<210> 17
<211> 54
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:nucleotide
      sequence of peptide 38 with in-frame stop codon
<400> 17
                                                                    54
actagtgggt tgctgaagct ggtgcaggct aagcgtaagt gttgtattag ttag
<210> 18
<211> 25
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence:peptide #40
<400> 18
Arg Trp Asp Pro Thr Arg Leu Leu Arg Phe Arg Phe Leu Arg Met Leu
                                      10
                  5
  1
Val Arg Arg Ser Arg Pro Val Arg Pro
             20
<210> 19
<211> 25
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence:peptide #88
<400> 19
Arg Leu Arg Arg Ile Cys Ser Gly Ile Leu Leu Ile Arg Arg Ile Leu
                  5
Gly Ile Phe Val Arg Pro Val Arg Pro
             20
<210> 20
<211> 40
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:variant of
      synthetic peptide #40 with N-terminus seven Lys
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and linker, K7\_40

<400> 20 Lys Lys Lys Lys Lys Lys Gly Glu Glu Ala Ala Lys Ala Arg Trp Asp Pro Thr Arg Leu Leu Arg Phe Arg Phe Leu Arg Met Leu Val Arg Arg Ser Arg Pro Val Arg Pro <210> 21 <211> 40 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence:variant of synthetic peptide #40 with N-terminus seven Lys and linker and alanine mutated residues, K7\_40 M <400> 21 Lys Lys Lys Lys Lys Lys Gly Glu Glu Ala Ala Lys Ala Arg 5 Trp Asp Pro Thr Arg Ala Leu Arg Ala Arg Phe Ala Arg Ala Leu Val 25 Arg Arg Ser Arg Pro Val Arg Pro 35 <210> 22 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: variant of synthetic peptide #41 with N-terminus seven Lys and linker, K7 41 <400> 22 Lys Lys Lys Lys Lys Lys Gly Glu Glu Ala Ala Lys Ala Gly Arg Gly Cys Ile Phe Arg Trp Arg Arg Gly Leu Arg Gly Met Met Arg

Leu Phe Lys

<210> 23

<211> 35

<212> PRT

<213> Artificial Sequence

35

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<220>
<223> Description of Artificial Sequence:variant of
      synthetic peptide #41 with N-terminus seven Lys
      and linker and alanine mutated residues, K7 41 M
<400> 23
Lys Lys Lys Lys Lys Lys Gly Glu Glu Ala Ala Lys Ala Gly
Arg Gly Cys Ile Phe Arg Ala Arg Gly Ala Arg Gly Met Ala Arg
                                 25
Ala Phe Lys
<210> 24
<211> 5
<212> PRT
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<223> Description of Artificial Sequence:5 arginine
      residues
<400> 24
Arg Arg Arg Arg
<210> 25
<211> 8
<212> PRT
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<400> 25
Gly Gly Glu Glu Ala Ala Lys Ala
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<210> 26
<211> 13
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<223> Description of Artificial Sequence: C-terminus of
      GFP and linker fused to biotinylated peptide 40,
      41 and 35
<400> 26
Gly Met Asp Glu Leu Tyr Lys Glu Glu Ala Ala Lys Ala
  1
                  5
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<210> 27
<211> 200
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 flexible linker
<220>
<221> MOD RES
<222> (6)..(200)
<223> Gly residues from position 6 to 200 may be present
 or absent
<400> 27
80
110
   100
130
       135
             140
165
          170
185
               190
Gly Gly Gly Gly Gly Gly
<210> 28
<211> 24
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9

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence:proliferating cell nuclear antigen (PCNA)-binding C-terminal peptide of tumor suppressor p21 (p21C) <400> 28 Lys Arg Arg Gln Thr Ser Met Thr Asp Phe Tyr His Ser, Lys Arg Arg 10 Leu Ile Phe Ser Lys Arg Lys Pro 20 <210> 29 <211> 25 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence:peptide #40 alanine mutant (M15A) <400> 29 Arg Trp Asp Pro Thr Arg Leu Leu Arg Phe Arg Phe Leu Arg Ala Leu 10 Val Arg Arg Ser Arg Pro Val Arg Pro 20 <210> 30 <211> 25 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence:peptide #40 alanine mutant (L13A/M15A) <400> 30 Arg Trp Asp Pro Thr Arg Leu Leu Arg Phe Arg Phe Ala Arg Ala Leu 10 Val Arg Arg Ser Arg Pro Val Arg Pro 20 <210> 31 <211> 25 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence:peptide #40 alanine mutant (F10A/L13A/M15A)

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<400> 31
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Val Arg Arg Ser Arg Pro Val Arg Pro
             20
<210> 32
<211> 20
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:peptide #41
      alanine mutant (L18A)
<400> 32
Gly Arg Gly Cys Ile Phe Arg Trp Arg Gly Leu Arg Gly Met Met
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Arg Ala Phe Lys
<210> 33
<211> 20
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<223> Description of Artificial Sequence:peptide #41
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<400> 33
Gly Arg Gly Cys Ile Phe Arg Trp Arg Arg Gly Leu Arg Gly Met Ala
                                     10
Arq Ala Phe Lys
<210> 34
<211> 20
<212> PRT
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<223> Description of Artificial Sequence:peptide #41
      alanine mutant (L12A/M16A/L18A)
<400> 34
Gly Arg Gly Cys Ile Phe Arg Trp Arg Gly Ala Arg Gly Met Ala
                                     10
Arg Ala Phe Lys
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20

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<210> 35
<211> 25
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<213> Artificial Sequence
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<400> 35
Arg Leu Arg Arg Ile Cys Ser Gly Ile Leu Leu Ile Arg Arg Ile Leu
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Gly Ala Phe Val Arg Pro Val Arg Pro
             20
<210> 36
<211> 25
<212> PRT
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<223> Description of Artificial Sequence:peptide #88
      alanine mutant (L16A/I18A)
<400> 36
Arg Leu Arg Arg Ile Cys Ser Gly Ile Leu Leu Ile Arg Arg Ile Ala
                                      10
Gly Ala Phe Val Arg Pro Val Arg Pro
             20
<210> 37
<211> 25
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:peptide #88
      alanine mutant (I12A/L16A/I18A)
<400> 37
Arg Leu Arg Arg Ile Cys Ser Gly Ile Leu Leu Ala Arg Arg Ile Ala
Gly Ala Phe Val Arg Pro Val Arg Pro
<210> 38
<211> 10
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<220>
<223> Description of Artificial Sequence:leucine-rich
      motif of HIV-1 Rev (amino acid position 75-84)
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<400> 38
Leu Pro Pro Leu Glu Arg Leu Thr Leu Asp
<210> 39
<211> 11
<212> PRT
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<223> Description of Artificial Sequence:leucine-rich
      motif of Mitogen-activated protein kinase kinase
      1, MAPKK (amino acid position 33-43)
<400> 39
Leu Gln Lys Lys Leu Glu Glu Leu Glu Leu Asp
                  5
<210> 40
<211> 12
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:leucine-rich
      motif of HTLV-1 Rex (amino acid position 82-93)
<400> 40
Leu Ser Ala Gln Leu Tyr Ser Ser Leu Ser Leu Asp
                  5
  1
                                      10
<210> 41
<211> 12
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:leucine-rich
      motif of Human homologue of mouse double minute
      2, Hdm-2 (amino acid position 190-200)
<400> 41
Ile Ser Leu Ser Phe Asp Glu Ser Leu Ala Leu Cys
<210> 42
<211> 11
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence:leucine-rich
      motif of Protein kinase inhibitor, PKI (amino acid
      position 38-48)
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<400> 42
Leu Ala Leu Lys Leu Ala Gly Leu Asp Ile Asn
<210> 43
<211> 17
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:C-terminal 17
      residues of cell division cycle 42 isoform 2,
      CDC42C (amino acid position 183-191)
<400> 43
Ala Ala Leu Glu Pro Pro Glu Thr Gln Pro Lys Arg Lys Cys Cys Ile
                                      10
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Phe
<210> 44
<211> 9
<212> PRT
<213> Artificial Sequence
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      #38Ndelta(1-8)
<400> 44
Gln Ala Lys Arg Lys Cys Cys Ile Ser
  1
<210> 45
<211> 4
<212> PRT
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<223> Description of Artificial Sequence:peptide
      #38Ndelta(1-13)
<400> 45
Cys Cys Ile Ser
  1
<210> 46
<211> 17
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      #38(T1A)
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<400> 46
Ala Ser Gly Leu Leu Lys Leu Val Gln Ala Lys Arg Lys Cys Cys Ile
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Ser
<210> 47
<211> 17
<212> PRT
<213> Artificial Sequence
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      #38 (S2A)
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Thr Ala Gly Leu Leu Lys Leu Val Gln Ala Lys Arg Lys Cys Cys Ile
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Ser
<210> 48
<211> 17
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      #38 (G3A)
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Thr Ser Ala Leu Leu Lys Leu Val Gln Ala Lys Arg Lys Cys Cys Ile
 1
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Ser
<210> 49
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      #38(L4A)
<400> 49
Thr Ser Gly Ala Leu Lys Leu Val Gln Ala Lys Arg Lys Cys Cys Ile
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Ser
<210> 50
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<220>
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      #38(L5A)
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Thr Ser Gly Leu Ala Lys Leu Val Gln Ala Lys Arg Lys Cys Cys Ile
Ser
<210> 51
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      #38 (K6A)
<400> 51
Thr Ser Gly Leu Leu Ala Leu Val Gln Ala Lys Arg Lys Cys Cys Ile
                                      10
Ser
<210> 52
<211> 17
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      #38(L7A)
<400> 52
Thr Ser Gly Leu Leu Lys Ala Val Gln Ala Lys Arg Lys Cys Cys Ile
 1
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Ser
<210> 53
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      #38 (V8A)
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Thr Ser Gly Leu Leu Lys Leu Ala Gln Ala Lys Arg Lys Cys Cys Ile
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Ser
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<210> 54
<211> 17
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      #38 (C14A)
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Thr Ser Gly Leu Leu Lys Leu Val Gln Ala Lys Arg Lys Ala Cys Ile
                                      10
Ser
<210> 55
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Arg Trp Asp Pro Thr Arg Leu Leu Arg Phe Arg Phe Leu Arg Met Leu
                  5
  1
                                      10
                                                           15
Val Arg Arg Ser Arg Pro Val Arg
             20
<210> 56
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<212> PRT
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<223> Description of Artificial Sequence: Human homologue
      of mouse double minute 2, HDM-2
Leu Ser Leu Ser Phe Asp Glu Ser Leu Ala Leu Cys
                  5
<210> 57
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<223> Description of Artificial Sequence:N-terminal
      sequence of peptides 35, 40 and 41 synthesized
      with C-terminus of GFP and spacer residues
<220>
<221> MOD_RES
<222> (1)
<223> Xaa = biotinylated Gly
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